



CGCTCAGGATACGACTTCCTCCCTAGACGATGGGATCCCGGGGATTATATAGCTCGATGATCT
TTCTCTATATCTCTCCGATCTGGGATATATACACACACACCTCCGGATAGCATGACTTCTT
CCCCCACT
CACACGATCT

PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy	OMIM	Books
Search <input type="text" value="Nucleotide"/> for <input type="text"/>							Go	Clear
Limits		Preview/Index		History		Clipboard		Details
Display	GenBank	Save	Text	Add to Clipboard				

☐ 1: AI732223. nf17g03.x5 NCI_CG...[gi:5053336] ProbeSet, Taxonomy, LinkOut

LOCUS AI732223 457 bp mRNA linear EST 13-DEC-1999
 DEFINITION nf17g03.x5 NCI_CGAP_Pr1 Homo sapiens cDNA clone IMAGE:914068
 similar to contains element PTR7 repetitive element mRNA
 sequence.
 ACCESSION AI732223
 VERSION AI732223.1 GI:5053336
 KEYWORDS EST.
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (bases 1 to 457)
 AUTHORS NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>.
 TITLE National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
 Tumor Gene Index
 JOURNAL Unpublished (1997)
 COMMENT Contact: Robert Strausberg, Ph.D.
 Email: cgapbs-r@mail.nih.gov
 Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
 Michael Emmert-Buck, M.D., Ph.D.
 cDNA Library Preparation: David B. Krizman, Ph.D.
 cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
 DNA Sequencing by: Washington University Genome Sequencing Center
 Clone distribution: NCI-CGAP clone distribution information can be
 found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

RECEIVED

APR 03 2002

TECH CENTER 1600/2900

This read is a RESEQUENCE of a previously sequenced human clone
 Original clone citation: National Cancer Institute, Cancer Genome
 Anatomy Project (CGAP), Tumor Gene Index
 This read has been verified (found to hit its original self in the
 correct orientation)
 Insert Length: 573 Std Error: 0.00
 Seq primer: -40UP from Gibco
 High quality sequence stop: 418.

FEATURES
 source Location/Qualifiers
 1..457
 /organism="Homo sapiens"
 /db_xref="taxon:9606"
 /clone="IMAGE:914068"
 /clone_lib="NCI_CGAP_Pr1"
 /sex="Male"
 /dev_stage="45 years old"
 /lab_host="DH10B"
 /note="Vector: pAMP10; Site_1: NotI; Site_2: EcoRI; 1st
 strand cDNA was primed with oligo(dT)17 on 50 ng of
 DNase-treated, total cellular RNA obtained from 5,000-10
 ,000 microdissected, histologically normal prostate
 epithelial cells. Double-stranded cDNA was ligated to
 EcoRI adaptors, 5 cycles of PCR applied to the cDNA with
 an adaptor-specific primer, and the resulting PCR product
 subcloned into pAMP10 by the UDG-cloning method (Life
 Technologies). Average insert size is 600 bp. NOTE: Not
 directionally cloned. This library was constructed by
 David Krizman."

BASE COUNT 92 a 135 c 105 g 125 t